



PCT09

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/831,307B

DATE: 01/27/2003
 TIME: 13:16:49

Input Set : A:\viraxseq.txt
 Output Set: N:\CRF4\01272003\I831307B.raw

4 <110> APPLICANT: The Macfarlane Burnet Centre for Medical Research Limited;
 6 Commonwealth Scientific and Industrial Research Organisation
 8 The Australian National University
 12 <120> TITLE OF INVENTION: Recombinant Viral Constructs and Methods Relating
 14 Thereto
 18 <130> FILE REFERENCE: 2231646/TDO
 C--> 22 <140> CURRENT APPLICATION NUMBER: US/09/831,307B
 C--> 24 <141> CURRENT FILING DATE: 2001-05-08
 28 <160> NUMBER OF SEQ ID NOS: 4
 32 <170> SOFTWARE: PatentIn Ver. 2.0

ERRORED SEQUENCES

84 <210> SEQ ID NO: 4
 86 <211> LENGTH: 37
 88 <212> TYPE: DNA
 90 <213> ORGANISM: mammalian
 94 <400> SEQUENCE: 4
 96 aaaggatcct ttagcttct taaaaaaaaa catatgg
 E--> 110 document2-08/11/01
 E--> 115 - 3 -

*Does Not Comply
Corrected Diskette Needed*

37

delete

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 01/27/2003
PATENT APPLICATION: US/09/831,307B TIME: 13:16:50

Input Set : A:\viraxseq.txt
Output Set: N:\CRF4\01272003\I831307B.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 2,4,6,8,10,12,14,16,18,20,22,24,26,28,30,32,34,36,38,40,42
Seq#:1; Line(s) 44,46,48,50,52
Seq#:2; Line(s) 54,56,58,60,62,64,66,68
Seq#:3; Line(s) 70,72,74,76,78,80,82,84
Seq#:4; Line(s) 86,88,90,92,94

VARIABLE LOCATION SUMMARY
PATENT APPLICATION: US/09/831,307B

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Input Set : A:\viraxseq.txt
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Use of n's or Xaa's(NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which
residue n or Xaa represents.

Seq#:4; N Pos. 42

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/831,307B

DATE: 01/27/2003
TIME: 13:16:50

Input Set : A:\viraxseq.txt
Output Set: N:\CRF4\01272003\I831307B.raw

L:22 M:270 C: Current Application Number differs, Replaced Application Number
L:24 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:110 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:4
L:110 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:48 SEQ:4
L:110 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10
L:110 M:112 C: (48) String data converted to lower case,
M:254 Repeated in SeqNo=4
L:115 M:252 E: No. of Seq. differs, <211> LENGTH:Input:37 Found:48 SEQ:4